

1
met ala gln trp glu met leu gln
ACTGC~~A~~CCCTAA~~T~~CAGAGCCCCAA ATG GCG CAG TGG GAA ATG CTG CAG

10
asn leu asp ser pro phe gln asp gln leu his gln leu tyr ser
AAT CTT GAC AGC CCC TTT CAG GAT CAG CTG CAC CAG CTT TAC TCG

20
his ser leu leu pro val asp ile arg gln tyr leu ala val trp
CAC AGC CTC CTG CCT GTG GAC ATT CGA CAG TAC TTG GCT GTC TGG

30
ile glu asp gln asn trp gln glu ala ala leu gly ser asp asp
ATT GAA GAC CAG AAC TGG CAG GAA GCT GCA CTT GGG AGT GAT GAT

40
ser lys ala thr met leu phe phe his phe leu asp gln leu asn.
TCC AAG GCT ACC ATG CTA TTC CAC TTC TTG GAT CAG CTG AAC

50
tyr glu cys gly arg cys ser gln asp pro glu ser leu leu leu
TAT GAG TGT GGC CGT TGC AGC CAG GAC CCA GAG TCC TTG TTG CTG

60
90
gln his asn leu arg lys phe cys arg asp ile gln pro phe ser
CAG CAC AAT TTG CCG AAA TTC TGC CCG GAC ATT CAG CCC TTT TCC

70
tyr glu cys gly arg cys ser gln asp pro glu ser leu leu leu
TAT GAG TGT GGC CGT TGC AGC CAG GAC CCA GAG TCC TTG TTG CTG

80
100
gln asp pro thr gln leu ala glu met ile phe asn leu leu leu
CAG GAT CCT ACC CAG TTG GCT GAG ATG ATC TTT AAC CTC CTT CTG

110
120
glu glu lys arg ile leu ile gln ala gln arg ala gln leu glu
GAA GAA AAA AGA ATT TTG ATC CAG GCT CAG AGG GCC CAA TTG GAA

130
140
gln gly glu pro val leu glu thr pro val glu ser gln gln his
CAA GGA GAG CCA GTT CTC GAA ACA CCT GTG GAG AGC CAG CAA CAT

150
glu ile glu ser arg ile leu asp leu arg ala met met glu lys
GAG ATT GAA TCC CGG ATC CTG GAT TTA AGG GCT ATG ATG GAG AAG

160
170
leu val lys ser ile ser gln leu lys asp gln gln asp val phe
CTG GTA AAA TCC ATC AGC CAA CTG AAA GAC CAG CAG GAT GTC TTC

180

Figure 1A

cys phe arg tyr lys ile gln ala lys gly lys thr pro ser leu
 TGC TTC CGA TAT AAG ATC CAG GCC AAA GGG AAG ACA CCC TCT CTG
 190
 asp pro his gln thr lys glu gln lys ile leu gln glu thr leu
 GAC CCC CAT CAG ACC AAA GAG CAG AAG ATT CTG CAG GAA ACT CTC
 200
 asn glu leu asp lys arg arg lys glu val leu asp ala ser lys
 AAT GAA CTG GAC AAA AGG AGA AAG GAG GTG CTG GAT GCC TCC AAA
 210
 ala leu leu gly arg leu thr thr leu ile glu leu leu leu pro
 GCA CTG CTA GGC CGA TTA ACT ACC CTA ATC GAG CTA CTG CTG CCA
 220
 lys leu glu glu trp lys ala gln gln lys ala cys ile arg
 AAG TTG GAG GAG TGG AAG GCC CAG CAA AAA GCC TGC ATC AGA
 230
 ala pro ile asp his gly leu glu gln leu glu thr trp phe thr
 GCT CCC ATT GAC CAC GGG TTG GAA CAG CTG GAG ACA TGG TTC ACA
 240
 ala gly ala lys leu leu phe his leu arg gln leu leu lys glu
 GCT GGA GCA AAG CTG TTG TTT CAC CTG AGG CAG CTG CTG AAG GAG
 250
 260
 leu lys gly leu ser cys leu val ser tyr gln asp asp pro leu
 CTG AAG GGA CTG AGT TGC CTG GTT AGC TAT CAG GAT GAC CCT CTG
 270
 280
 thr lys gly val asp leu arg asn ala gln val thr glu leu leu
 ACC AAA GGG GTG GAC CTA CGC AAC GCC CAG GTC ACA GAG TTG CTA
 290
 300
 gln arg leu leu his arg ala phe val val glu thr gln pro cys
 CAG CGT CTG CTC CAC AGA GCC TTT GTG GTC GAA ACC CAG CCC TGC
 310
 320
 met pro gln thr pro his arg pro leu ile leu lys thr gly ser
 ATG CCC CAA ACT CCC CAT CGA CCC CTC ATC CTC AAG ACT GGC AGC
 330
 340
 lys phe thr val arg thr arg leu leu val arg leu gln glu gly
 AAG TTC ACC GTC CGA ACA AAG CTG CTG GTG AGA CTC CAG GAA GGC
 350
 360
 asn glu ser leu thr val glu val ser ile asp arg asn pro pro
 AAT GAG TCA CTG ACT GTG GAA GTC TCC ATT GAC AGG AAT CCT CCT
 370
 380
 gln leu gln gly phe arg lys phe asn ile leu thr ser asn gln
 CAA TTA CAA GGC TTC CGG AAG TTC AAC ATT CTG ACT TCA AAC CAG
 390
 lys thr leu thr pro glu lys gly gln ser gln gly leu ile trp

Figure 1B

AAA ACT TTG ACC CCC GAG AAG GGG CAG AGT CAG GGT TTG ATT TGG
 400
 asp phe gly tyr leu thr leu val glu gln arg ser gly gly ser
 GAC TTT GGT TAC CTG ACT CTG GTG GAG CAA CGT TCA GGT TCA
 420
 gly lys gly ser asn lys gly pro leu gly val thr glu glu leu
 GGA AAG GGC AGC ATAT AAG GGG CCA CTA GGT GTG ACA GAG GAA CTG
 430
 his ile ile ser phe thr val lys tyr thr tyr gln gly leu lys
 CAC ATC ATC AGC TTC ACG GTC AAA TAT ACC TAC CAG GGT CTG AAG
 450
 gln glu leu lys thr asp thr leu pro val val ile ile ser asn
 CAG GAG CTG AAA ACG GAC ACC CTC CCT GTG GTG ATT ATT TCC AAC
 460
 met asn gln leu ser ile ala trp ala ser val leu trp phe asn
 ATG AAC CAG CTC TCA ATT GCC TGG GCT TCA GTT CTC TGG TTC ATT
 480
 leu leu ser pro asn leu gln asn gln gln phe phe ser asn pro
 TTG CTC AGC CCA AAC CTT CAG AAC CAG CAG TTC TCC AAC CCC
 490
 pro lys ala pro trp ser leu leu gly pro ala leu ser trp gln
 CCC AAG GCC CCC TGG AGC TTG CTG GGC CCT GCT CTC AGT TGG CAG
 510
 phe ser ser tyr val gly arg gly leu asn ser asp gln leu ser
 TTC TCC TCC TAT GTT GGC CGA GGC CTC AAC TCA GAC CAG CTG AGC
 520
 met leu arg asn lys leu phe gly gln asn cys arg thr glu asp
 ATG CTG AGA AAC AAG CTG TTC GGG CAG AAC TGT AGG ACT GAG GAT
 540
 pro leu leu ser trp ala asp phe thr lys arg glu ser pro pro
 CCA TTA TTG TCC TGG GCT GAC TTC ACT AAC CGA GAG AGC CCT CCT
 550
 gly lys leu pro phe trp thr trp leu asp lys ile leu glu leu
 GGC AAG TTA CCA TTC TGG AGC TGG CTG GAC AAA ATT CTG GAG TTG
 570
 val his asp his leu lys asp leu trp asn asp gly arg ile met
 GTA CAT GAC CAC CTG AAG GAT CTC TGG ATT GAT GGA CGC ATC ATG
 580
 gly phe val ser arg ser gln glu arg arg leu leu lys lys thr
 GGC TTT GTG AGT CGG AGC CAG GAG CGC CGG CTG CTG AAG AAC ACC
 600
 met ser gly thr phe leu leu arg phe ser glu ser ser glu gly
 ATG TCT GGC ACC TTT CTA CTG CGC TTC AGT GAA TCG TCA GAA GGG

Figure 1C

600-1-195 A

(Sheet 4 of 42)

610
 gly ile thr cys ser trp val glu his gln asp asp asp lys val
 GGC ATT ACC TGC TCC TGG GTG GAG CAC CAG GAT GAT GAC AAG CTG
 630
 leu ile tyr ser val gln pro tyr thr lys glu val leu gln ser
 CTC ATC TAC TCT GTG CAA CCG TAC ACG AAG GAG GTG CTG CAG TCA
 640
 650
 leu pro leu thr glu ile ile arg his tyr gln leu leu thr glu
 CTC CCG CTG ACT GAA ATC ATC CGC CAT TAC CAG TTG CTC ACT GAG
 660
 glu asn ile pro glu asn pro leu arg phe leu tyr pro arg ile
 GAG AAT ATA CCT GAA AAC CCA CTG CGC TTC CTC TAT CCC CGA ATC
 670
 680
 pro arg asp glu ala phe gly cys tyr tyr gln glu lys val asn
 CCC CGG GAT GAA GCT TTT GGG TGC TAC TAC CAG GAG AAA GTT AAT
 690
 leu gln glu arg arg lys tyr leu lys his arg leu ile val val
 CTC CAG GAA CGG AAG AAA TAC CTG AAA CAC AGG CTC ATT GTG GTC
 700
 710
 ser asn arg gln val asp glu leu gln gln pro leu glu leu lys
 TCT AAT AGA CAG GTG GAT GAA CTG CAA CCG CTG GAG CTT AAG
 720
 730
 pro glu pro glu leu glu ser leu glu leu glu leu gly leu val
 CCA GAG CCA GAG CTG GAG TCA TTA GAG CTG GAA CTA GGG CTG GTC
 pro glu pro glu leu ser leu asp leu glu pro leu leu lys ala
 CCA GAG CCA GAG CTC AGC CTG GAC TTA GAG CCA CTG CTG AAG GCA
 750
 gly leu asp leu gly pro glu leu glu ser val leu glu ser th
 GGG CTG GAT CTG GGG CCA GAG CTA GAG TCT GTG CTG GAG TCC ACT
 760
 770
 leu glu pro val ile glu pro thr leu cys met val ser gln th
 CTG GAG CCT GTG ATA GAG CCC ACA CTA TGC ATG GTA TCA CAA AC
 780
 val pro glu pro asp gln gly pro val ser gln pro val pro gl
 GTG CCA GAG CCA GAC CAA GGA CCT GTA TCA CAG CCA GTG CCA GA
 790
 800
 pro asp leu pro cys asp leu arg his leu asn thr glu pro me
 CCA GAT TTG CCC TGT GAT CTG AGA CAT TGT AAC ACT GAG CCA AT
 810
 glu ile phe arg asn cys val lys ile glu glu ile met pro as
 GAA ATC TTC AGA AAC TGT GTA AAG ATT GAA GAA ATC ATG CCG AA

Figure 1D

020
gly asp pro leu leu ala gly gln asn thr val asp glu val tyr
GGT GAC CCA CTG TTG GCT GGC CAG AAC ACC GTG GAT GAG GTT TAC

030

val ser arg pro ser his phe tyr thr asp gly pro-leu met pro
GTC TCC CGC CCC AGC CAC TTC TAC ACT GAT GGA CCC TTG ATG CCT

040

ser asp phe AM
TCT GAC TTC TAG GAAACCACATTCCCTCTGTTCTTTCATATCTCTTTGCCCTTCATA
CTCCTCATAGCATGATATTGTTCTCCAGGATGGAAATCAGGCATGTGTCCCTTCAGC
TGTGTTAACTGTTCAAAACTCAGGCCCTGTGTGACTCCATTGGGGTGAGAGGTGAAGCATA
ACATGGGTACAGAGGGGACAAACAAATGAATCAGAACAGATGCTGAGCCATAGGTCTAAATA
GGATCCTGGAGGCTCCCTGCTGTGCTGGGAGGTATAGGGGTCTGGGGGAGGCCAGGGC
AGTTGACAGGTACTTGGAGGGCTCAGGGCAGTGGCTTCTT'TCCAGTATGGAAGGATTTC
ACATTTTAATAGTTGGTTAGGCTAAACTGGTCATACTGGCATTGGCCTTGGTGGGGAGC
ACAGACACAGGATAGGACTCCATTCTTCTTCCATTCCATTCTCATGTCTAGGATAACTTGC
TTTCTTCTTCTTACTCCTGGCTCAAGCCCTGAAATTCTTCTTCTTGCAGGGTTG
AGAGCTTCTGCCTTAGCCTACCATGTGAAACTCTACCCCTGAAAGAAAAGGATGGATAGGA
AGTAGACCTCTTTCTT'TACCAAGTC'CCTCCCCTACTCTGCCCTAAGCTGGCTGTACC
TGTTCCCTCCCCATAAATGATCCTGCCAAATCTAAGAAAAGAAA

Figure 1E

ATTAACCTCGCCGAGCCCCCTCCGCAGACTCTGGCCGGAAAGTTTATTTGTATGCCATCCTCGA
 GAGCTGTCTAGGTTAACGTTGCACCTGTGTATATAACCTCGACAGCTTGGCACCTAACGTGCTGTGCG
 TAGCTGCTCCCTTGTTGAATCCCCAGGCCCTGTTGGGGACAAGGTGGCAGG ATG TCT CAG TGG Met Ser Gln Trp
 Tyr Glu Leu Gln Gln Leu Asp Ser Lys Phe Leu Glu Gln Val His Gln Leu Tyr
 TAC GAA CTT CAG CAG CTT GAC TCA AAA TTC CTG GAG CAG GTT CAC CAG CTT TAT
 Asp Asp Ser Phe Pro Met Glu Ile Arg Gln Tyr Leu Ala Gln Trp Leu Glu Lys
 GAT GAC AGT TTT CCC ATG GAA ATC AGA CAG TAC CTG GCA CAG TGG TTA GAA AAG
 Gln Asp Trp Glu His Ala Ala Asn Asp Val Ser Phe Ala Thr Ile Arg Phe His
 CAA GAC TGG GAG CAC GCT GCC AAT GAT GTT TCA TTT GCC ACC ATC CGT TTT CAT
 Asp Leu Leu Ser Gln Leu Asp Gln Tyr Ser Arg Phe Ser Leu Glu Asn Asn
 GAC CTC CTG TCA CAG CTG GAT GAT CAA TAT AGT CGC TTT TCT TTG GAG AAT AAC
 Phe Leu Leu Gln His Asn Ile Arg Lys Ser Lys Arg Asn Leu Gln Asp Asn Phe
 TTC TTG CTA CAG CAT AAC ATA AGG AAA AGC AAG CGT AAT CTT CAG GAT AAT TTT
 Gln Glu Asp Pro Ile Gln Met Ser Met Ile Ile Tyr Ser Cys Leu Lys Glu Glu
 CAG GAA GAC CCA ATC CAG ATG TCT ATC ATT TAC AGC TGT CTG AAG GAA GAA
 Arg Lys Ile Leu Glu Asn Ala Gln Arg Phe Asn Gln Ala Gln Ser Gly Asn Ile
 AGG AAA ATT CTG GAA AAC GCC CAG AGA TTT AAT CAG GCT CAG TCG GGG AAT ATT
 Gln Ser Thr Val Met Leu Asp Lys Gln Lys Glu Leu Asp Ser Lys Val Arg Asn
 CAG AGC ACA GTG ATG TTA GAC AAA CAG GAG CTT GAC AGT AAA GTC AGA AAT
 Val Lys Asp Lys Val Met Cys Ile Glu His Glu Ile Lys Ser Leu Glu Asp Leu
 GTG AAG GAC AAG GTT ATG TGT ATA GAG CAT GAA ATC AAG AGC CTG GAA GAT TTA
 Gln Asp Glu Tyr Asp Phe Lys Cys Lys Thr Leu Gln Asn Arg Glu His Glu Thr
 CAA GAT GAA TAT GAC TTC AAA TGC AAA ACC TTG CAG AAC AGA GAA CAC GAG ACC
 Asn Gly Val Ala Lys Ser Asp Gln Lys Gln Glu Gln Leu Leu Lys Lys Met
 AAT GGT GTG GCA AAG AGT GAT CAG AAA CAA GAA CAG CTG TTA CTC AAG AAG ATG
 Tyr Leu Met Leu Asp Asn Lys Arg Lys Glu Val Val His Lys Ile Ile Glu Leu
 TAT TTA ATG CTT GAC AAT AAG AGA AAG GAA GTA GTT CAC AAA ATA ATA GAG TTG
 Leu Asn Val Thr Glu Leu Thr Gln Asn Ala Leu Ile Asn Asp Glu Leu Val Glu
 CTG AAT GTC ACT GAA CTT ACC CAG AAT GCC CTG ATT AAT GAT GAA CTA GTG GAG
 Trp Lys Arg Arg Gln Gln Ser Ala Cys Ile Gly Gly Pro Pro Asn Ala Cys Leu
 TGG AAG CGG AGA CAG AGC GCC TGT ATT GGG GGG CCG CCC AAT GCT TGC TTG
 Asp Gln Leu Gln Asn Trp Phe Thr Ile Val Ala Glu Ser Leu Gln Gln Val Arg
 GAT CAG CTG CAG AAC TGG TTC ACT ATA GTT GCG GAG AGT CTG CAG CAA GTT CGG
 Gln Gln Leu Lys Lys Leu Glu Glu Leu Glu Gln Lys Tyr Thr Tyr Glu His Asp
 CAG CAG CTT AAA AAG TTG GAG GAA TTG GAA CAG AAA TAC ACC TAC GAA CAT GAC
 Pro Ile Thr Lys Asn Lys Gln Val Leu Trp Asp Arg Thr Phe Ser Leu Phe Gln
 CCT ATC ACA AAA AAC AAA CAA GTG TTA TGG GAC CGC ACC TTC AGT CTT TTC CAG

Figure 2A

Gln Leu Ile Gln Ser Ser Phe Val Val Glu Arg Gln Pro Cys Met Pro Thr His
 CAG CTC ATT CAG AGC TCG TTT GTG GTG GAA AGA CAG CCC TGC ATG CCA ACG CAC

 Pro Gln Arg Pro Leu Val Leu Lys Thr Gly Val Phe Thr Val Lys Leu Arg
 CCT CAG AGG CCG CTG GTC TTG AAG ACA GGG GTC CAG TTC ACT GTG AAG TTG AGA

 Leu Leu Val Lys Leu Gln Glu Leu Asn Tyr Asn Leu Lys Val Lys Val Leu Phe
 CTG TTG GTG AAA TTG CAA GAG CTG AAT TAT AAT TTG AAA GTC AAA GTC TTA TTT

 Asp Lys Asp Val Asn Glu Arg Asn Thr Val Lys Gly Phe Arg Lys Phe Asn Ile
 GAT AAA GAT GTG AAT GAG AGA AAT ACA GTA AAA GGA TTT AGG AAG TTC AAC ATT

 Leu Gly Thr His Thr Lys Val Met Asn Met Glu Glu Ser Thr Asn Gly Ser Leu
 TTG GGC ACG CAC ACA AAA GTG ATG AAC ATG GAG GAG TCC ACC AAT GGC AGT CTG

 Ala Ala Glu Phe Arg His Leu Gln Leu Lys Glu Gln Lys Asn Ala Gly Thr Arg
 GCG GCT GAA TTT CGG CAC CTG CAA TTG AAA GAA CAG AAA AAT GCT GGC ACC AGA

 Thr Asn Glu Gly Pro Leu Ile Val Thr Glu Glu Leu His Ser Leu Ser Phe Glu
 ACG AAT GAG GGT CCT CTC ATC GTT ACT GAA GAG CTT CAC TCC CTT AGT TTT GAA

 Thr Gln Leu Cys Gln Pro Gly Leu Val Ile Asp Leu Glu Thr Thr Ser Leu Pro
 ACC CAA TTG TGC CAG CCT GGT TTG GTA ATT GAC CTC GAG ACG ACC TCT CTG CCC

 Val Val Val Ile Ser Asn Val Ser Gln Leu Pro Ser Gly Trp Ala Ser Ile Leu
 GTT GTG GTG ATC TCC AAC GTC AGC CAG CTC CCG AGC GGT TGG GCC TCC ATC CTT

 Trp Tyr Asn Met Leu Val Ala Glu Pro Arg Asn Leu Ser Phe Phe Leu Thr Pro
 TGG TAC AAC ATG CTG GTG GCG GAA CCC AGG AAT CTG TCC TTC TTC CTG ACT CCA

 Pro Cys Ala Arg Trp Ala Gln Leu Ser Glu Val Leu Ser Trp Gln Phe Ser Ser
 CCA TGT GCA CGA TGG GCT CAG CTT TCA GAA GTG CTG AGT TGG CAG TTT TCT TCT

 Val Thr Lys Arg Gly Leu Asn Val Asp Gln Leu Asn Met Leu Gly Glu Lys Leu
 GTC ACC AAA AGA GGT CTC AAT GTG GAC CAG AAC ATG TTG GGA GAG AAG CTT

 Leu Gly Pro Asn Ala Ser Pro Asp Gly Leu Ile Pro Trp Thr Arg Phe Cys Lys
 CTT GGT CCT AAC GCC AGC CCC GAT GGT CTC ATT CCG TGG ATT GAA AGC ATC CTA

 Glu Asn Ile Asn Asp Lys Asn Phe Pro Phe Trp Leu Trp Ile Glu Ser Ile Leu
 GAA AAT ATA AAT GAT AAA AAT TTT CCC TTC TGG CTT TGG ATT GAA AGC ATC CTA

 Glu Leu Ile Lys Lys His Leu Leu Pro Leu Trp Asn Asp Gly Cys Ile Met Gly
 GAA CTC ATT AAA AAA CAC CTG CTC CCT CTC TGG AAT GAT GGG TGC ATC ATG GGC

 Phe Ile Ser Lys Glu Arg Glu Arg Ala Leu Leu Lys Asp Gln Gln Pro Gly Thr
 TTC ATC AGC AAG GAG CGA GAG CGT GCC CTG TTG AAG GAC CAG CAG CCG GGG ACC

 Phe Leu Leu Arg Phe Ser Glu Ser Ser Arg Glu Gly Ala Ile Thr Phe Thr Trp
 TTC CTG CGG TTC AGT GAG AGC TCC CGG GAA GGG GCC ATC ACA TTC ACA TGG

 Val Glu Arg Ser Gln Asn Gly Glu Pro Asp Phe His Ala Val Glu Pro Tyr
 GTG GAG CGG TCC CAG AAC GGA GGC GAA CCT GAC TTC CAT GCG GTT GAA CCC TAC

 Thr Lys Lys Glu Leu Ser Ala Val Thr Phe Pro Asp Ile Ile Arg Asn Tyr Lys
 ACG AAG AAA GAA CTT TCT GCT GTT ACT TTC CCT GAC ATC ATT CGC AAT TAC AAA

 Val Met Ala Ala Glu Asn Ile Pro Glu Asn Pro Leu Lys Tyr Leu Tyr Pro Asn
 GTC ATG GCT GCT GAG AAT ATT CCT GAG AAT CCC CTG AAG TAT CTG TAT CCA AAT

Figure 2B

Ile Asp Lys Asp His Ala Phe Gly Lys Tyr Ser Arg Pro Lys Glu Ala Pro
ATT GAC AAA GAC CAT GCC TTT GGA AAG TAT TAC TCC AGG CCA AAG GAA GCA CCA

Glu Pro Met Glu Leu Asp Gly Pro Lys Gly Thr Gly Tyr Ile Lys Thr Glu Leu
GAG CCA ATG GAA CTT GAT GGC CCT AAA GGA ACT GGA TAT ATC AAG ACT GAG TTG

Ile Ser Val Ser Glu Val His Pro Ser Arg Leu Gln Thr Thr Asp Asn Leu Leu
ATT TCT GTG TCT GAA GTT CAC CCT TCT AGA CTT CAG ACC ACA GAC AAC CTG CTC

Pro Met Ser Pro Glu Glu Phe Asp Glu Val Ser Arg Ile Val Gly Ser Val Glu
CCC ATG TCT CCT GAG GAG TTT GAC GAG GTG TCT CGG ATA GTG GGC TCT GTA GAA

Phe Asp Ser Met Met Asn Thr Val
TTC GAC AGT ATG AAC ACA GTA TAGAGCATGAATTTTTCATCTTCTCTGGCGACAGTTT

TCCTTCTCATCTGTGATTCCCTCCTGCTACTCTGTTCTCACATCCTGTGTTCTAGGAAATGAAAGAA

AGGCCAGCAAATTCGCTGCAACCTGTTGATAGCAAGTGAATTTCTCTAACTCAGAAACATCAGTTACTC

TGAAGGGCATCATGCATCTTACTGAAGGTAAAATTGAAAGCATTCTCTGAAGAGTGGGTTACAAGTGA

AAAACATCCAGATAACCCAAAGTATCAGGACGAGAATGAGGGCTTGGAAAGGAGAAGTTAACCAAC

ATCTAGCAAATGTTATGCATAAAGTCAGTGCCCACTGTTAGGGTGGATAATCAGTGGTTATTTA

GGGAAGTGTGCTTAGGAACGGTAAATTCTGTGGAGAATTCTACATGTTCTTGCTTAAGTGT

AACTGGCAGTTCCATTGGTTACCTGTGAAATAGTCAGGCAAGTTATATAAATTATCAGTCC

TCTTCAAGGTAGCCATCATGGATCTGGTAGGGGAAAATGTGTATTTATTACATCTTCACATTGGCT

ATTTAAAGACAAAGACAAATTCTGTTCTGAGAAGAGAATATTAGCTTACTGTTGTTAGGCTTAATG

ACACTAGCTAATCAATAGAAGGATGTACATTCCAATTACAAGTTGTGATATCCAAGCTGAA

TACATTCTGCTTCACTTGGTCACATACAATTATTTACAGTTCTCCAAGGGAGTTAGGCTATTCA

ACCACTCATTCAAAGTTGAAATTAAACCATAGATGTAGATAAACTCAGAAATTAAATTCAATTGTTAAA

TGGGCTACTTGTCTTTGTTAGGGTGGTATTAGTCTATTAGCCACAAAATTGGGAAAGGAGTAG

AAAAAGCAGTAACGTACAACCTGAATAATACACCAGAGATAATGAGAATCAGATCATTCAAAACTCAT

TTCCCTATGTAACGTGCAATTGAGAACTGCATATGTTGCTGATATATGTGTTTCACTTGCAGATGGTT

Figure 2C

CCATTCTCTCCTGTACTTTTCCAGACACTTTTGAGTGGATGATGTTCGTGAAGTATACTGTATT
TTACCTTTTCTTCCTTATCACTGACACACAAAAGTAGATTAAGAGATGGGTTGACAAGGTTCTCCCTT
TTACATACTGCTGTCTATGTGGCTGTATCTGTTTCCACTACTGCTACCACAACTATATTATCATGCAA
ATGCTGTATTCTTCTTGGTGGAGATAAAGATTCTTGAGTTTGTAAATTAAAGCTAAAGTATCTG
TATTGCATTAATATAATATCGACACAGTGCTTCCGTGGCACTGCATACAACTGAGGCCCTCTCTCA
GTTTTATATAGATGGCGAGAACCTAAGTTCAAGTTGATTTACAATTGAAATGACTAAAAACAAAGAAG
ACAAACATTAACAAATATTGTTCTA

Figure 2D

ATTAACCTCTGCCGAGCCCTCCGACACTCTGCCGGAAAGTTCATTTGCTGTATGCCATCCTCGA
 GAGCTGCTAGGTTAACGTCGCACTCTGTGTATATAACCTCGACAGTCTTGCACCTAACGTGCTGTGCG
 Met Ser Gln Trp
 TAGCTGCTCCTTGGTTGAATCCCCAGGCCCTGTTGGGCACAAGGTGGCAGG ATG TCT CAG TGG
 Tyr Glu Leu Gln Gln Leu Asp Ser Lys Phe Leu Glu Gln Val His Gln Leu Tyr
 TAC GAA CTT CAG CAG CTT GAC TCA AAA TTC CTG GAG CAG GTT CAC CAG CTT TAT
 Asp Asp Ser Phe Pro Met Glu Ile Arg Gln Tyr Leu Ala Gln Trp Leu Glu Lys
 GAT GAC AGT TTT CCC ATG GAA ATC AGA CAG TAC CTG GCA CAG TGG TTA GAA AAG
 Gln Asp Trp Glu His Ala Ala Asn Asp Val Ser Phe Ala Thr Ile Arg Phe His
 CAA GAC TGG GAG CAC GCT GCC AAT GAT GTT TCA TTT GCC ACC ATC CGT TTT CAT
 Asp Leu Leu Ser Gln Leu Asp Asp Gln Tyr Ser Arg Phe Ser Leu Glu Asn Asn
 GAC CTC CTG TCA CAG CTG GAT GAT CAA TAT AGT CGC TTT TCT TTG GAG AAT AAC
 Phe Leu Leu Gln His Asn Ile Arg Lys Ser Lys Arg Asn Leu Gln Asp Asn Phe
 TTC TTG CTA CAG CAT AAC ATA AGG AAA AGC AAG CGT AAT CTT CAG GAT AAT TTT
 Gln Glu Asp Pro Ile Gln Met Ser Met Ile Ile Tyr Ser Cys Leu Lys Glu Glu
 CAG GAA GAC CCA ATC CAG ATG TCT ATG ATC ATT TAC AGC TGT CTG AAG GAA GAA
 Arg Lys Ile Leu Glu Asn Ala Gln Arg Phe Asn Gln Ala Gln Ser Gly Asn Ile
 AGG AAA ATT CTG GAA AAC GCC CAG AGA TTT AAT CAG GCT CAG TCG GGG AAT ATT
 Gln Ser Thr Val Met Leu Asp Lys Gln Lys Glu Leu Asp Ser Lys Val Arg Asn
 CAG AGC ACA GTG ATG TTA GAC AAA CAG GAG CTT GAC AGT AAA GTC AGA AAT
 Val Lys Asp Lys Val Met Cys Ile Glu His Glu Ile Lys Ser Leu Glu Asp Leu
 GTG AAG GAC AAG GTT ATG TGT ATA GAG CAT GAA ATC AAG AGC CTG GAA GAT TTA
 Gln Asp Glu Tyr Asp Phe Lys Cys Lys Thr Leu Gln Asn Arg Glu His Glu Thr
 CAA GAT GAA TAT GAC TTC AAA TGC AAA ACC TTG CAG AAC AGA GAA CAC GAG ACC
 Asn Gly Val Ala Lys Ser Asp Gln Lys Gln Glu Gln Leu Leu Lys Lys Met
 AAT GGT GTG GCA AAG AGT GAT CAG AAA CAA GAA CAG CTG TTA CTC AAG AAG ATG
 Tyr Leu Met Leu Asp Asn Lys Arg Lys Glu Val Val His Lys Ile Ile Glu Leu
 TAT TTA ATG CTT GAC AAT AAG AGA AAG GAA GTA GTT CAC AAA ATA ATA GAG TTG
 Leu Asn Val Thr Glu Leu Thr Gln Asn Ala Leu Ile Asn Asp Glu Leu Val Glu
 CTG AAT GTC ACT GAA CTT ACC CAG AAT GCC CTG ATT AAT GAT GAA CTA GTG GAG
 Trp Lys Arg Arg Gln Gln Ser Ala Cys Ile Gly Gly Pro Pro Asn Ala Cys Leu
 TGG AAG CGG AGA CAG CAG AGC GCC TGT ATT GGG GGG CCG CCC AAT GCT TGC TTG
 Asp Gln Leu Gln Asn Trp Phe Thr Ile Val Ala Glu Ser Leu Gln Gln Val Arg
 GAT CAG CTG CAG AAC TGG TTC ACT ATA GTT GCG GAG AGT CTG CAG CAA GTT CGG
 Gln Gln Leu Lys Lys Leu Glu Glu Leu Glu Gln Lys Tyr Thr Tyr Glu His Asp
 CAG CAG CTT AAA AAG TTG GAG GAA TTG GAA CAG AAA TAC ACC TAC GAA CAT GAC
 Pro Ile Thr Lys Asn Lys Gln Val Leu Trp Asp Arg Thr Phe Ser Leu Phe Gln
 CCT ATC ACA AAA AAC CAA GTG TTA TGG GAC CGC ACC TTC AGT CTT TTC CAG

Figure 3A

Gln Leu Ile Gln Ser Ser Phe Val Val Glu Arg Gln Pro Cys Met Pro Thr His
 CAG CTC ATT CAG AGC TCG TTT GTG GTG GAA AGA CAG CCC TGC ATG CCA ACG CAC

 Pro Gln Arg Pro Leu Val Leu Lys Thr Gly Val Gln Phe Thr Val Lys Leu Arg
 CCT CAG AGG CCG CTG GTC TTG AAG ACA GGG GTC CAG TTC ACT GTG AAG TTG AGA

 Leu Leu Val Lys Leu Gln Glu Leu Asn Tyr Asn Leu Lys Val Lys Val Leu Phe
 CTG TTG GTG AAA TTG CAA GAG CTG AAT TAT TTG AAA GTC AAA GTC TTA TTT

 Asp Lys Asp Val Asn Glu Arg Asn Thr Val Lys Gly Phe Arg Lys Phe Asn Ile
 GAT AAA GAT GTG AAT GAG AGA AAT ACA GTA AAA GGA TTT AGG AAG TTC AAC ATT

 Leu Gly Thr His Thr Lys Val Met Asn Met Glu Glu Ser Thr Asn Gly Ser Leu
 TTG GGC ACG CAC ACA AAA GTG ATG AAC ATG GAG GAG TCC ACC AAT GGC AGT CTG

 Ala Ala Glu Phe Arg His Leu Gln Leu Lys Glu Gln Lys Asn Ala Gly Thr Arg
 GCG GCT GAA TTT CGG CAC CTG CAA TTG AAA GAA CAG AAA AAT GCT GGC ACC AGA

 Thr Asn Glu Gly Pro Leu Ile Val Thr Glu Glu Leu His Ser Leu Ser Phe Glu
 ACG AAT GAG GGT CCT CTC ATC GTT ACT GAA GAG CTT CAC TCC CTT AGT TTT GAA

 Thr Gln Leu Cys Gln Pro Gly Leu Val Ile Asp Leu Glu Thr Thr Ser Leu Pro
 ACC CAA TTG TGC CAG CCT GGT TTG GTA ATT GAC CTC GAG ACG ACC TCT CTG CCC

 Val Val Val Ile Ser Asn Val Ser Gln Leu Pro Ser Gly Trp Ala Ser Ile Leu
 GTT GTG ATC TCC AAC GTC AGC CAG CTC CCG AGC GGT TGG GCC TCC ATC CTT

 Trp Tyr Asn Met Leu Val Ala Glu Pro Arg Asn Leu Ser Phe Phe Leu Thr Pro
 TGG TAC AAC ATG CTG GTG GCG GAA CCC AGG AAT CTG TCC TTC TCC CTG ACT CCA

 Pro Cys Ala Arg Trp Ala Gln Leu Ser Glu Val Leu Ser Trp Gln Phe Ser Ser
 CCA TGT GCA CGA TGG GCT CAG CTT TCA GAA GTG CTG AGT TGG CAG TTT TCT TCT

 Val Thr Lys Arg Gly Leu Asn Val Asp Gln Leu Asn Met Leu Gly Glu Lys Leu
 GTC ACC AAA AGA GGT CTC AAT GTG GAC CAG CTG AAC ATG TTG GGA GAG AAG CTT

 Leu Gly Pro Asn Ala Ser Pro Asp Gly Leu Ile Pro Trp Thr Arg Phe Cys Lys
 CTT GGT CCT AAC GCC AGC CCC GAT GGT CTC ATT CCG TGG ACG AGG TTT TGT AAG

 Glu Asn Ile Asn Asp Lys Asn Phe Pro Phe Trp Leu Trp Ile Glu Ser Ile Leu
 GAA AAT ATA AAT GAT AAA AAT TTT CCC TTC TGG CTT TGG ATT GAA AGC ATC CTA

 Glu Leu Ile Lys Lys His Leu Leu Pro Leu Trp Asn Asp Gly Cys Ile Met Gly
 GAA CTC ATT AAA AAA CAC CTG CTC CCT CTC TGG AAT GAT GGG TGC ATC ATG GGC

 Phe Ile Ser Lys Glu Arg Glu Arg Ala Leu Leu Lys Asp Gln Gln Pro Gly Thr
 TTC ATC AGC AAG GAG CGA GAG CGT GCC CTG TTG AAG GAC CAG CAG CCG GGG ACC

 Phe Leu Leu Arg Phe Ser Glu Ser Ser Arg Glu Gly Ala Ile Thr Phe Thr Trp
 TTC CTG CTG CGG TTC AGT GAG AGC TCC CGG GAA GGG GCC ATC ACA TTC ACA TGG

 Val Glu Arg Ser Gln Asn Gly Gly Glu Pro Asp Phe His Ala Val Glu Pro Tyr
 GTG GAG CGG TCC CAG AAC GGA GGC GAA CCT GAC TTC CAT GCG GTT GAA CCC TAC

 Thr Lys Lys Glu Leu Ser Ala Val Thr Phe Pro Asp Ile Ile Arg Asn Tyr Lys
 ACG AAG AAA GAA CTT TCT GCT GTT ACT TTC CCT GAC ATC ATT CGC AAT TAC AAA

 Val Met Ala Ala Glu Asn Ile Pro Glu Asn Pro Leu Lys Tyr Leu Tyr Pro Asn
 GTC ATG GCT GGT GAG AAT ATT CCT GAG AAT CCC CTG AAG TAT CTG TAT CCA AAT

Figure 3B

600-1-195 A

(Sheet 12 of 42)

Ile Asp Lys Asp His Ala Phe Gly Lys Tyr Tyr Ser Arg Pro Lys Glu Ala Pro
ATT GAC AAA GAC CAT GCC TTT GGA AAG TAT TAC TCC AGG CCA AAG GAA GCA CCA

Glu Pro Met Glu Leu Asp Gly Pro Lys Gly Thr Gly Tyr Ile Lys Thr Glu Leu
GAG CCA ATG GAA CTT GAT GGC CCT AAA GGA ACT GGA TAT ATC AAG ACT GAG TTG

Ile Ser Val Ser Glu Val
ATT TCT GTG TCT GAA GTG TAAGTGAACACAGAAGAGTGACATGTTACAAACCTCAAGCCAGCCT

TGCTCCTGGCTGGGGCTGTTGAAGATGCTTGATTTTACTTTCCATTGTAATTGCTATGCCATCACAG

CTGAACCTGTTGAGATCCCCGTGTTACTGCCTATCAGCATTACTACTTTAAAAAAAAGCCA

AAAACCAAATTGTATTTAAGGTATATAAATTTCACAAACTGATACCCTTGAAAAAGTATAAATAAA

TGAGCAAAAGTTGAA

Figure 3C

600-1-195 A

(Sheet 13 of 42)

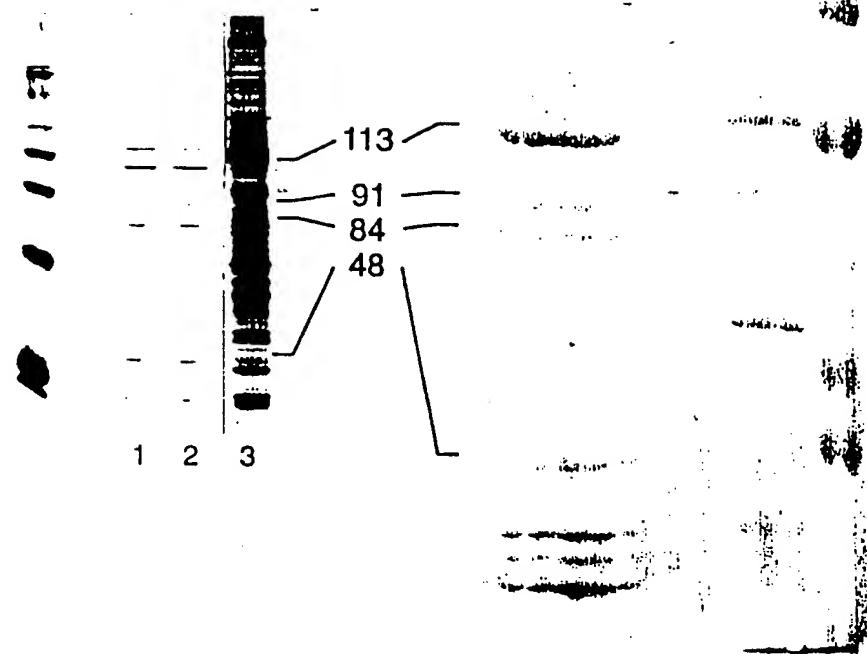


Figure 4

600-1-195 A

(Sheet 14 of 42)

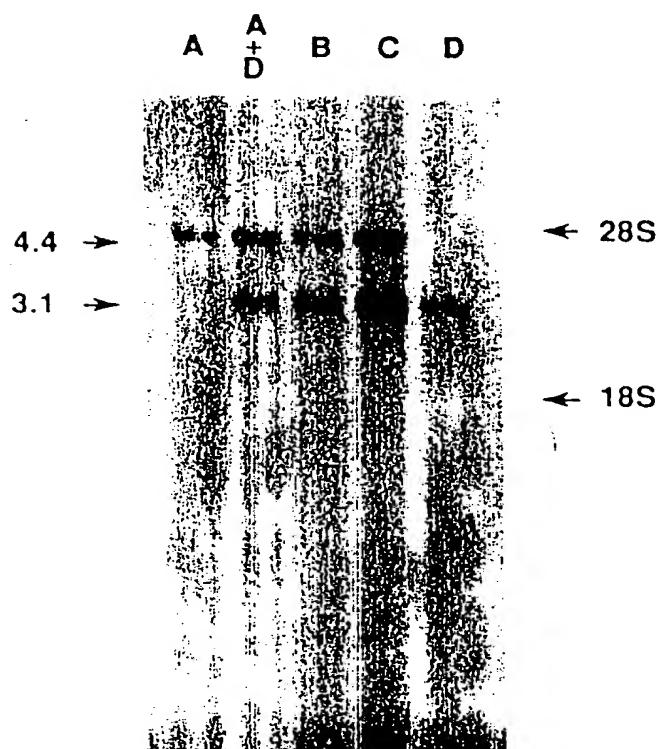
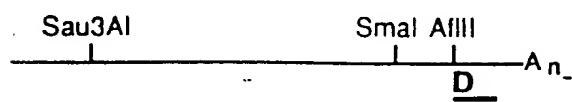
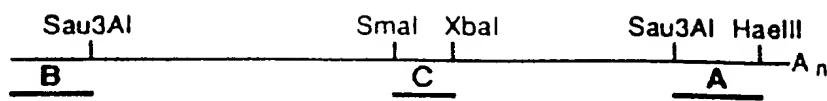


Figure 5

1 MSQWYELOQLDSKFLEQVHQLYDDSFPMETRQYLAQWLEKQDWHEAANDV
51 SFATIRFIIDLLSQLDDQYSRFSENNFLLQHINIRKSKRNLQDNFQEDP1Q
101 MSHIIYSCLKEERKILENAQRFNQAQSGNIQSTVMILDQKELDSKVRNVK
151 DKVMCIEHEIKSLEDLQDEYDFKCKTLQNREHETNGVAKSDQKQEQLLK
201 KTYLMLDNKRKEVVIIKKIIIELNVTELTONALINDELVEWKRRQQSACIGG
251 PPNACLDQLQQVROQLKKLEELEQKYTYEIIDPITKKNQVLWDRTFSLFQQ
301 LIQSSFVVERQPCMPTIIPQRPLVLTGVQFTVKLRLLVKLQELNYNLKVK
351 VLFDKDVNERNTVKGFRKFNI LGTH¹²⁷ KVMNMEESTNGSLAAEFRHLQLKE
401 QKHNAGTRTNEGPLIVTEELIISLSFETQLCOPGLVIDLETTSLPVVVISNV
451 SQLPSGWASILWYNMLVAEPRNLSFFLTTPCARWAQLSEVLSWQFSSVTK
501 RGLNYDOLNMLGEKLLGPNASPDGLIPWTRFCKENINDKNFPFWLWIESI
119
551 LELIKKHLPLWNNDGCIMGFISKERERALLKDQQPGTELLRFSESSREGA
601 ITFTWVERSQNGGEPDFIAVEPYTKKELS AVTFPDIIRNYKVMAAENIPE
113a
651 NPLKYLYPNLKDIAFGKYYSRPKEAPEPHEDGPKGTYIKTELISVSE
113b
701 VHP SRLQT TDNL LPMSPEEFDEVSRIVGSVEFDMMNTV
↑
last amino acid of 84 kd

Figure 6

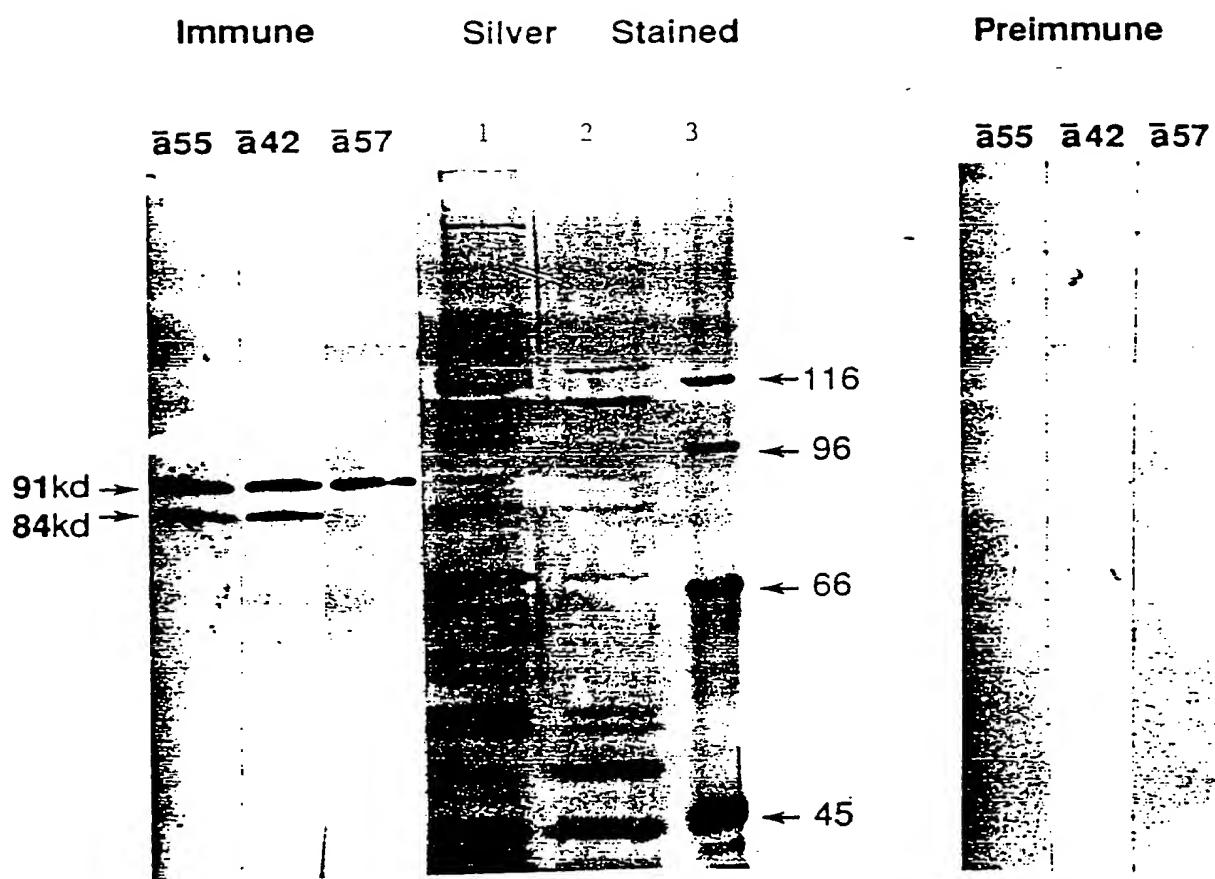


Figure 7A

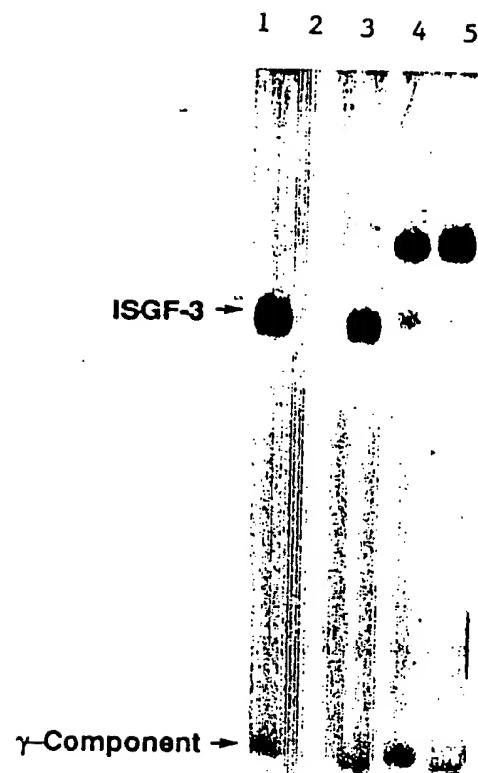


Figure 7B

1: MAQWEMLQNLDSPFQDQLHQLYSHSLLPVDIRQYLAVWIEDQNWQEAAALGSDDSKATMLF
 61: FHFLDQLNYECGRCSQDPESLLLQHNLRKFCRDIQPFSQDPTQLAEMIFNLLLEEKRILI
 121: QAQRAQLEQGEPVLET PVE SQQHEIESRILDLRAMMEKLVKSISQLKDQQDVECFRYKIQ
 181: AKGKTPSLDPHQTKEQKILQETLNELDKRRKEVLDASKALLGRLTTIELLLPKLEEWKA
 241: QQQKACIRAPIDHGLEQLETWETAGAKLLEHLRQLLKELKGLSCLVSYQDDPLTKGVDLR
 301: NAQVTELLQRLLHRAFVVETQPCMPQTPHRPLILKTGSKFTVRTRLLVRLQEGNESLTVE
 361: VSIDRNPPQLQGERKENILTSNQKTLTPEKGQSQGLIWIWDFGYLTVEQRSGGSGKGSNK
 421: PLGVTEELHIISFTVKYTYYQGLKQELKTDTLPVVIISNMNQLSIAWASVLWFNLLSPNLQ
 481: NQQFFSNPPKAPWSLLGPALSQFSSYVGRGLNSDQLSMLRNKLFQNCRTEDPLLSWAD
 541: FTKRESPPGKLPFWTWLDKILELVHDHLKDLWNDGRIMGFVSRSQERRLLKKTMSGTFLL
 601: RFSSESSEGGITCSWVEHQDDDKVLIYSVQPYTKEVLQSLPLTEIIRHYOLLTEENIPENP
 661: LRFLYPRIPRDEAFGCYYQEKVNLQERRKYLKHRLIVVSNRQVDELOQPPLKPBPSLES
 721: L E L E L G L V P E P E L S L D L E P L L K A G L D L G P E L E S V L E S T L E P V I E P T L C M V S Q T V P E P D Q G
 781: P V S Q P V P B P D L P C D L R H L N T B P M B I F R N C V K I B B I M P N G D P L L A G Q N T V D E V Y V S R P S H F
 841: YTDGPLMPSDF

Figure 8 A

113 kDa [MAQWEMILQLDSDPFDQDCHOLYSHSLUPVDIROYLAIVWIEDONMQEALGSDDSKATMLF]
 91/84 kDa [MSQWYETQOLDSKELEGVHOLYDDSFPMIEIROYLAQWLEKQDMHEHAN--NDVSFAIRF]

61 FHFIDQDNYECCRGSQDPESELQLQHNLRKFCRDICP-FSQDPTOLAEMIFNLLKEEKRL
 57 HQLLSQUDDOYSRFSLE-NNFLQLQHNLRKSKRNLDQEQEDPIQMSMIIYSCLEKEERKL

120 ICAQRAQLEGGEPVLETIPVESQHQHEIESRIILDRAHMHEKLVKSIISQIKDQQDVFCFRYK-
 117 ENAQNFNQAGSGNIQSITVMDKOKELDSKVRNVDRKVMCIEHEIKSLEDQDEYDERCKT

179 IQAKGKTPS--LDPHOTKECKILQETLNEIDKRKEVLDASKAELGRITTLIE--ULLPK
 177 LQNREHETNGVAKSDOKQECOLLKKMYLMLQNKREEVHKIIEIL-NVTELTQNALINDE

235 LEEWKAQOQKACIRAPIDHGLEQLETHFTAGAKLLFHRLQULKEKGGLSCLVSYQDDPLT
 236 LVEWKRRQOQSACIGGPPNACIDQLOQ-----QVROQQLKMEELEQKYTYEHDBIW

295 KGVDLRNAQVTETLORILHRNFVVEIOPCMQPTPHRPLILKTKSKFTVRTRLLVRLQEGN
 285 KNKQVLWDRTFSLFOQLIQSSFVVEROPCMPTIIOPRLVVLKTVQVFTVKURLLVKGELIN

355 ESUTVEVSIQRNPPQ---LQGFRKFNIITSNOKTLTPEKGQSQGLIWDFGYLTILVEQRSG
 345 YNUKVKVLFQDKDVNERNTVKGFRKFNIIGTHITKVHNMEESTNGSLAAERHQLKEQKNA

412 GSGKGSNWKSPLGVTTEELHIIISFTVKYTYQGLKQEILKTDTLPVVVISNMNQLSIAHASVILW
 405 GT--RTNEGPLIVTEELHISLISFETQLCQPGIVIDLETTSLPVVVISNVSQLPSGHASVILW

472 FNLLSPNLQOFFSNPRKAPMSLUGPALSQFSSYVCRGLNSDOLSMLRNKLFCONCRT
 463 YNMIVAEPRNLSEFLTPPCARMACUSEVLSHOESSVTKRGVLNVDQLNHLGEKILCPNASP

532 EDPILSWADFTKRESPPGKLIPFMWTWLDKILELVHDILKDLWNDRIMGFVSRSCERRLLK
 523 DG-LIPWTRECKENINDKNFPFMWIESILELIKILLPLWNDCIMGFISKERERALLK

592 KTMSTGTFLLRFSESS-EGGITCSWVEH-QDDDKVLIYSMOPYTKEVLOSLPLTEIIRHKO
 582 DQQPCTFLLRFSESSREGAIIFTWVERSONGGEPDFIIAMEPYTKKEELSAVTFPDILIRNKK

650 LLTEENIPENPLRFUYPPRIPRDEAFCCYY-----QEKVNQERR--KMLKIRLIMVSNR
 642 VMALENIPENPIKYLYPNIOKDIIAFGKYYSRPKEAPEPMELDGPKGTGMIKTELISVSEV

702 QVDELQOQPLELKP
 702 HPSRLQTTDNLYLP

Figure 8B

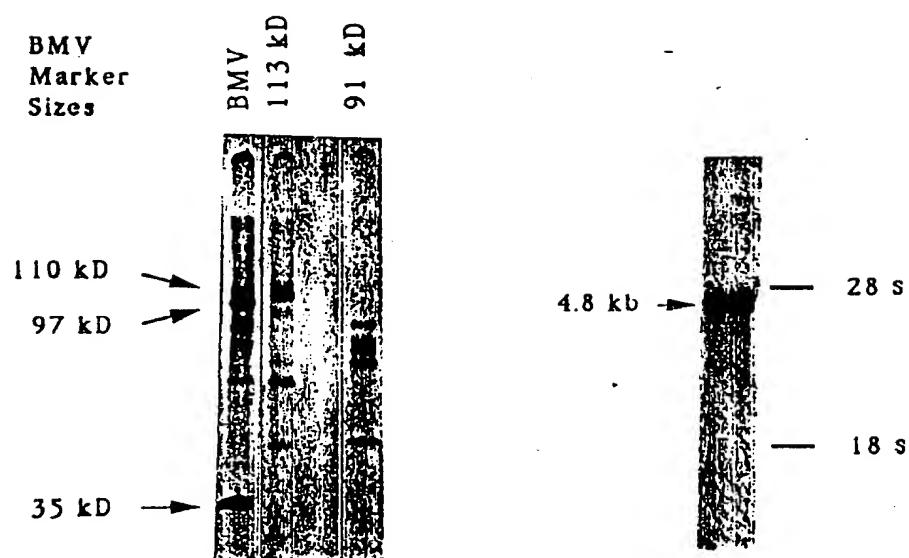


Figure 9

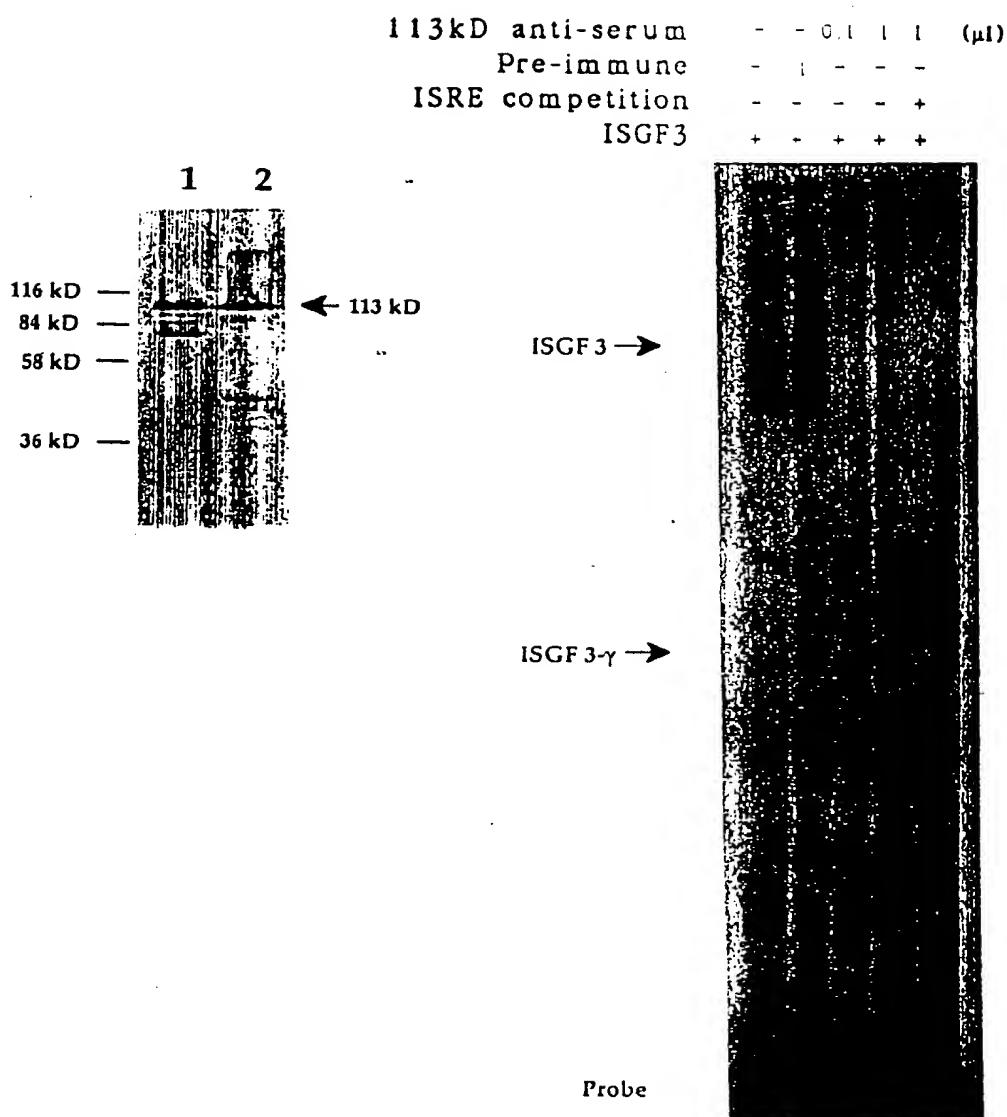
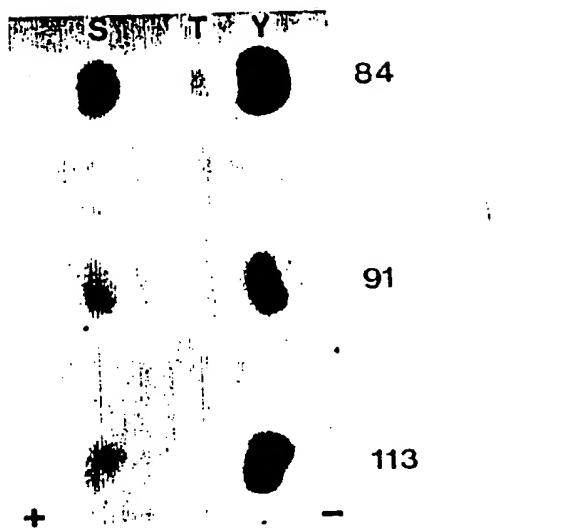


Figure 10

Figure 11

1 2 3 4 5 6 7

**Figure 12****Figure 11, 12**

1 MSQWFELQQQL DSKFLEQVHQ LYDDSFPMEI RQYLAQWLEK QDWEHAAYDV
51 SFATIRFHDL LSQLEDDQYSR FSLENNFLLQ HNIRKSKRNL QDNFQEDPVQ
101 MSMIIYNCLK EERKILENAQ RFNQAQEGNI QNTVMLDKQK ELDISKVRNVK
151 DQVMCIEQEI KTLEELQDEY DFKCKTSQNR EGEANGVAKS DQKQEQLLLH
201 KMFLMLDNKR KEIIHKIREL LNSIELTQNT LINDELVEWK RRQQSACIGG
251 PPNACLDQLQ TWFTIVAEYL QQIRQQLKKL EELEQKFTEP PDPITKNKQV
301 LSDRTFLLFQ QLIQSSFVVE RQPCMPTHPQ RPLVLKTGVQ FTVKSRLLVK
351 LQESNLLTKV KCHFDKDVE KNTVKGFRKF NILGTHTKVM NMEESTNGSL
401 AAELRHLQLK EQKNAGNRTN EGPLIVTEEL HSLSFETQLC QPGLVIDLET
451 TSLPVVVVISN VSQLPSGWAS ILWYNMLVTE PRNLSFFLNP PCAWWSQLSE
501 VLSWQFSSVT KRGLNADQLS MLGEKLLGPN AGPDGLIPWT RFCKENINDK
551 NESEFWPWIDT ILELIKNDLL CLWNDGCIMG FISKEERERAL LKDQQPGTFL
601 LRFSESSREG AITFTWVERS QNGGEPDFHA VEPTYKKELS AVTFPDIIRN
651 YKVMAAENIP ENPLKYLYPN IDKDHAFGKY YSRPKEAPEP MELDDPKRTG
701 YIKTELISVS EVHPSRLQTT DNLLPMSPEE FDEMSRIVGP EFDSMMSTV

Figure 13A

1 caggatgtca cagtggttcg agcttcagca gctggactcc aagttcctgg
51 agcaggtcca ccagctgtac gatgacagt tccccatgga aatcagacag
101 tacctggccc agtggctgga aaagcaagac tgggagcacg ctgcctatga
151 tgtctcgaaa gcgaccatcc gettccatga cctccctcta cagctggacg
201 accagtacag ccgctttct ctggagaata atttcttgtt gcagcacaac
251 atacggaaaa gcaagcgtaa tctccaggat aacttccaag aagatcccg
301 acagatgtcc atgatcatct acaactgtct gaaggaagaa aggaagattt
351 tggaaaatgc ccaaagattt aatcaggccc aggaggggaaa tattcagaac
401 actgtgtatgt tagataaaca gaaggagctg gacagtaaag tcagaaatgt
451 gaaggatcaa gtcatgtgca tagagcagga aatcaagacc ctagaagaat
501 tacaagatga atatgacttt aaatgcaaaa cctctcagaa cagagaaggt
551 gaagccaatg gtgtggcgaa gagcgaccaa aaacaggaac agctgctgct
601 ccacaagatg ttttaatgc ttgacaataa gagaaggag ataattcaca
651 aaatcagaga gttgctgaat tccatcgagc tcactcagaa cactctgatt
701 aatgacgagc tcgtggagtg gaagcgaagg cagcagagcg cctgcacatcg
751 gggaccgccc aacgcctgcc tggatcagct gcaaacgtgg ttcaccattg
801 ttgcagagac cctgcagcag atccgtcagc agctaaaaa gctggaggag
851 ttggaacaga aattcaccta tgagcccgac cctattacaa aaaacaagca
901 ggtgttgtca gatcgaacct tcctccctt ccagcagctc attcagagct
951 ctttcgtggt agaacgcacag ccgtgcacatgc ccactcaccc gcagaggccc
1001 ctggcttgc agactgggtt acagttcaact gtcacgtcga gactgttgt
1051 gaaattgc当地 gatcgaatc tattaacgaa agtggaaatgt cacttgtaca
1101 aagatgtgaa cgagaaaaac acagttaaag gatggggaa gttcaacatc
1151 ttgggtacgc acacaaaagt gatgaacatg gaaaaatcca ccaacggaaag
1201 tctggcagct gagctccgac acctgcaact gaaggaacag aaaaacgctg
1251 ggaacagaac taatgagggg cctctcattt gtcacggaga acttcactct
1301 ctttagctttt ggacccagtt gtgcacggca ggctgggtga ttgacctgg
1351 gaccacctct cttcctgtcg tggtgatctc caacgtcagc cagctccccca

Figure 13B

1401 gtggctggc gtctatcctg tggtaacaaca tgctggtgac agagcccagg
1451 aatctctcct tttccctgaa ccccccggtgc gcgtgggtt cccagctctc
1501 agaggtgtg agttggcagt ttcatcaagt caccaagaga ggtctgaacg
1551 cagaccagct gagcatgctg ggagagaagc tgctgggcc taatgctggc
1601 cctgatggtc ttattccatg gacaaggaaa tgtaaggaaa atattaatga
1651 taaaaatttc tccttctggc cttggattga caccatccta gagctcatTA
1701 agaacgacct gctgtgcctc tggaatgatg ggtgcattat gggcttcata
1751 agcaaggagc gagaacgcgc tctgctcaag gaccagcagc cagggacggt
1801 cctgcttaga ttcatgtaga gctcccgaa agggggccatc acattcacat
1851 gggtggaaacg gtcccagaac ggaggtgaac ctgacttcca tgccgtggag
1901 ccctacacga aaaaagaact ttcatgtttt acattccag atattattcg
1951 caactacaaa gtcatggctg ccgagaacat accagagaat cccctgaagt
2001 atctgtaccc caatattgac aaagaccacg cctttggaa gtattattcc
2051 agaccaaagg aagcaccaga accgatggag cttgacgacc ctaagcgaac
2101 tggataacatc aagactgagt tgatttctgt gtctgaagtc cacccttcta
2151 gacttcagac cacagacaac ctgcttcca tgtctccaga ggagtttgat
2201 gagatgtccc ggatagtggg ccccgaattt gacagtatga tgagcacagt
2251 ataaacacga atttctctc ggcgaca

Figure 13C

1 MSQWNQVQQL EIKFLEQVDQ FYDDNFPMEI RHLLAQWIET QDWEVASNNE
51 TMATILLQNL LIQLDEQLGR VSKEKNLLLI HNLKRIRKVL QGKFHGNPMH
101 VAVVISNCLR EERRILAAAN MPIQGPLEKS LQSSSVSERQ RNVEHKVSAI
151 KNSVQMTEQD TKYLEDLQDE FDYRYKTIQT MDQGDKNSIL VNQEVLTLQ
201 EMLNSLDFKR KEALSKMTQI VNETDLLMNS MLLEELQDWK KRRIACIGG
251 PLHNGLDQLQ NCFTLLAESL FQLRQQLEKL QEQSTKMTYE GDPIPAQRah
301 LLERATFLIY NLFKNSEVVE RHACMPTHPQ RPMVLKTLIQ FTVKLRLLIK
351 LPELNYQVKV KASIDKNVST LSNRRFVLCG THVKAMSSEE SSNGSLSVEL
401 DIATQGDEVQ YWSKGNEGCH MVTTEELHSIT FETQICLYGL TINLETSSLp
451 VVMISNVSQL PNAWASIIWY NVSTNDSQL VFFNNPPSVT LGQLLEVMSW
501 QFSSYVGRGL NSEQLNMLAE KLTVQSNYND GHLTWAKFCK EHLPGKTFTF
551 WTWLEAIDL IKKHILPLWI DGYIMGFVSK EKERLLIKDK MPGTFLLRFs
601 ESHLGGITFT WVDQSENGEV RFHSVEPYNK GRSLALAFAD ILRDYKVIMA
651 ENIPENPLKY LYPDIPKDKA FGKHYSSQPC EVSRPTERGD KGYPSPVFIP
701 ISTIRSDSTE PQSPSDLPPM SPSAYAVLRE NLSPTTIETA MNSPYSAE

Figure 14A

1 tgccactacc tggacggaga gagagagagc agcatgtctc agtggaatca
51 agtccaacaa ttagaaatca agtttttggg gcaaggatagat cagttctatg
101 atgacaacct tcctatggaa atccggcattc tgctagctca gtggatttag
151 actcaagact gggaaatggc ttctaacaat gaaactatgg caacaattct
201 gcttc当地 aacttaafac aattggatga acagttgggg cgggtttcca
251 aagaaaaaaaaa tctgctattt attcacaatc taaagagaat tagaaaagtt
301 ct当地 caggca agtttcatgg aaatccaatg catgttagctg tggtaatttc
351 aaattgctta agggaaagaga ggagaatatt ggctgcagcc aacatgccta
401 tccaggacc tctggagaaa tc当地 tacaga gttcttcagt ttctgaaaga
451 caaaggaatg tggacaccaa agtgtctgcc attaaaaaca gtgtgcagat
501 gacagaacaa gataccaaat acttagaaga cctgcaagat gagtttact
551 acaggtaaa aacaattcag acaatggatc agggtgacaa aaacagtatc
601 ct当地 gaacc aggaagttt gacactgctg caagaaatgc ttaatagtct
651 ggacttcaag agaaaggaag cactcagtaa gatgacgcag atagtgaacg
701 agacagacct gctcatgaac agcatgcttc tagaagagct gcaggactgg
751 aaaaagcggc acaggattgc ctgcattggg ggcccgtcc acaatggct
801 ggaccagctt cagaactgct ttaccctact ggcagagat cttttccaac
851 tc当地 acagca actggagaaa ctacaggagc aatctactaa aatgacctat

Figure 14B

901 gaaggggatc ccatccctgc tcaaagagca cacccctgg aaagagctac
951 cttcctgatc tacaacctt tcaagaactc atttgtggtc gagcgacacg
1001 catgcattgcc aacgcaccct cagaggccga tggtacttaa aaccctcatt
1051 cagttcactg taaaactgag attactaata aaattgcgg aactaaacta
1101 tcaggtgaaa gtaaaggcggt ccattgacaa gaatgttca actctaagca
1151 atagaagatt tgtgctttgt ggaactcacg tcaaagctat gtccagttag
1201 gaatcttcca atgggagcct ctcaagtggag tttagacattt caacccaagg
1251 agatgaagtg cagtaactgga gtaaaggaaa cgagggctgc cacatggta
1301 cagaggagtt gcattccata acctttgaga cccagatctg cctctatggc
1351 ctcaccattta accttagagac cagctcatta cctgtcgtga tgatttctaa
1401 tgtcagccaa ctacctaattt catgggcattc catcatggg tacaatgtat
1451 caactaacga ctcccagaac ttgggtttct ttaataaccc tccatctgtc
1501 actttggggcc aactccctgga agtgatgagc tggcaatttt catcctatgt
1551 cggtcgtggc ctttaattcag agcagctcaa catgctggca gagaagctca
1601 cagttcagtc taactacaat gatggtcacc tcacccctggc caagttctgc
1651 aaggaacatt tgcctggcaa aacatttacc ttctggactt ggcttgaagc
1701 aatattggac ctaataaaaa aacatattct tccccctctgg attgtatgggt
1751 acatcatggg atttgttagt aaagagaagg aacgggttct gctcaaagat
1801 aaaatgcctg ggacatttt gttaagattc agtgagagcc atcttggagg

Figure 14C

1851 gataaccttc acctgggtgg accaatctga aaatggagaa gtgagattcc
1901 actctgtaga accctacaac aaaggagac tgtcggctct ggcccttcgct
1951 gacatcctgc gagactacaa gtttatcatg gctgaaaaca tccctgaaaa
2001 ccctctgaag tacctctacc ctgacattcc caaagacaaa gccttggca
2051 aacactacag ctcccagccg tgcgaaatctt caagaccaac cgaacgggaa
2101 gacaagggtt acgtccccctc tgttttatc cccatttcaa caatccgaag
2151 cgattccacg gagccacaat ctccattcaga ctttctcccc atgtctccaa
2201 gtgcatatgc tgtgctgaga gaaaacctga gcccaacgac aattgaaaact
2251 gcaatgaatt cccatattc tgctaatga cggtgcaaac ggacacttta
2301 aagaaggaag cagatgaaac tggagagtgt tctttaccat agatcacaat
2351 ttatttcttc ggcttgtaa atacc

Figure 14D

1 MAQWNQLOQL DTRYLKQLHQ LYSDFPMEL RQFLAPWIES QDWAYAASKE
51 SHATL VFHNL LGEIDQQYSR FLQESNVLYQ HNLRIKQFL QSRYLEKPME
101 IARIVARCLW EESRLLQTAA TAAQQGGQAN HPTAAVVTEK QQMLEQHLQD
151 VRKRVQDLEQ KMKVVENLQD DFDFNYKTLK SQGDMQDLNG NNQSVTRQKM
201 QQLEQMLTAL DQMRRSIVSE LAGLLSAMEY VQKTLTDEEL ADWKRRPEIA
251 CIGGPPNICL DRLENWITSL AESQLQTRQQ IKKLEELQQK VSYKGDPIVQ
301 HRPMLEERIV ELFRNLMKSA FVVERQPCMP MHPDRPLVIK TGVQFTTKVR
351 LLVKFPELNY QLKIKVCIDK DSGDVAALRG SRKFNILGTN TKVMNMEESN
401 NGSLSAEFKH LTLREQRCGN GGRANCDASL IVTEELHLIT FETEVYHQGL
451 KIDLETHSLP VVVISNICQM PNAWASILWY NMLTNNPKNV NFFTKPPIGT
501 WDQVAEVL SW QFSSTTKRGL SIEQLTTLAE KLLGPGVNYS GCQITWAKEFC
551 KENMAGKGFS FWVWLDNIID LVKKYILALW NEGYIMGFIS KERERAILST
601 KPPGTFLRF SESSKEGGVT FTWVEKDISG KTQIQSVEPY TKQQLNNMSF
651 AEIIMGYKIM DATNILVSPL VYLYPDIPKE EAEGKYCRPE SQEHPEADPG
701 SAAPYLKTF ICVTPPTCSN TIDLPMSPRT LDSLMQFGNN GEGAEP SAGG
751 QFESLTFDMD LTSECATSPM

Figure 15A

1 gccgcgacca gccaggccgg ccagtcgggc tcagccccga gacagtcgag
51 acccctgact gcagcaggat ggctcagtgg aaccagctgc agcagctgga
101 cacacgctac ctgaaggcgc tgcaccagct gtacagcgac acgttccccca
151 tggagctgcg gcagttcctg gcaccttggg ttgagagtca agactggca
201 tatgcagcca gcaaagagtc acatgccacg ttggtgtttc ataatcttt
251 gggtaaaatt gaccagcaat atagccgatt cctgcaagag tccaatgtcc
301 tctatcagca caaccttcga agaatcaagc agtttctgca gagcaggtat
351 cttgagaagc caatggaaat tgccggatc gtggcccgat gcctgtggga
401 agagtctcgc ctccctccaga cggcagccac ggcagcccag caagggggcc
451 aggccaacca cccaacagcc gccgtagtga cagagaagca gcagatgttg
501 gagcagcatc ttcaaggatgt ccggaagcga gtgcaggatc tagaacagaa
551 aatgaaggtg gtggagaacc tccaggacga ctttgatttc aactacaaaa
601 ccctcaagag ccaaggagac atgcaggatc tgaatggaaa caaccagtct
651 gtgaccagac agaagatgca gcagctggaa cagatgctca cagccctgga
701 ccagatgcgg agaagcattt tgagttagt ggcggggctc ttgtcagcaa
751 tggagtacgt gcagaagaca ctgactgtat aagagctggc tgactggaag
801 aggcggccag agatcgcggtt catcgaggc cctcccaaca tctgcctgga
851 ccgtctggaa aactggataa cttcattagc agaatctcaa cttcagaccc

Figure 15B

901 gccaacaaat taagaaaactg gaggagctgc agcagaaaagt gtcctacaag
951 ggcgacccta tcgtgcagca ccggcccatg ctggaggaga ggatcggtga
1001 gctgttcaga aacttaatga agagtgcctt cgtggtgag cgccagccct
1051 gcatgccat gcacccggac cggcccttag tcatcaagac tggtgtccag
1101 ttaccacga aagtcaaggtt gctggtaaaa tttcctgagt tgaattatca
1151 gcttaaaatt aaagtgttca ttgataaaga ctctgggat gttgctgccc
1201 tcagagggtc tcggaaattt aacattctgg gcacgaacac aaaagtgtatg
1251 aacatggagg agtctaacaa cggcagcctg tctgcagagt tcaagcacct
1301 gacccttagg gagcagagat gtggaatgg aggccgtgcc aattgtgtatg
1351 cctcctttagt cgtgactgag gagctgcacc tgatcacctt cgagactgag
1401 gtgtaccacc aaggcctcaa gattgaccta gagacccact cttgccagt
1451 tgtggtgatc tccaacatct gtcagatgcc aaatgcttgg gcatcaatcc
1501 tgtggtataa catgctgacc aataacccca agaacgtgaa cttttcact
1551 aagccgccaa ttgaaacctg ggaccaagtg gccgagggtgc tcagctggca
1601 gttctcgatcc accaccaagc gagggtctgag catcgagcag ctgacaacgc
1651 tggctgagaa gtccttaggg cctgggtgtga actactcagg gtgtcagatc
1701 acatgggcta aattctgcaa agaaaacatg gctggcaagg gcttcctt
1751 ctgggtctgg ctagacaata tcatcgaccc tggaaaaag tataatcttgg
1801 cccttggaa tgaagggtac atcatgggtt tcatcagcaa ggagcgggag

Figure 15C

1851 cgggccatcc taagcacaaa gccccgggc acttcctac tgcgcttcag
1901 cgagagcagc aaagaaggag gggtaacttt cacttgggtg gaaaaggaca
1951 tcagtggcaa gacccagatc cagtcgttag agccatacac caagcagcag
2001 ctgaacaaca tgtcatttgc tgaatcatc atgggctata agatcatgga
2051 tgcgaccaac atcctggtgt ctccacttgt ctacctctac cccgacattc
2101 ccaaggagga ggcatttgg aagtactgta ggcccggag ccaggagcac
2151 cccgaagccg acccaggttag tgctgcccc tacctgaaga ccaagttcat
2201 ctgtgtgaca ccaacgacct gcagcaatac cattgacctg ccatgtccc
2251 cccgcacttt agattcattt atgcagttt gaaataacgg tgaaggtgct
2301 gagccctcag caggagggca gttttagtcg ctcacgtttt acatggatct
2351 gacctcgag tgtgctacct ccccatgtg aggagctgaa accagaagct
2401 gcagagacgt gacttgagac acctgccccg tgctccaccc ctaagcagcc
2451 gaacccata tcgtctgaaa ctcctaactt tgggttcca gatTTTTT
2501 tttaatttcc tacttctgct atctttgggc aatctggca ctTTTTaaaa
2551 gagagaaatg agtgagtgtg ggtgataaac tgttatgtaa agaggagaga
2601 cctctgagtc tggggatggg gctgagagca gaagggaggc aaaggggaac
2651 acctcctgtc ctgccccctt gccctcctt ttcagcagct cgggggttgg
2701 ttgttagaca agtgccctt ggtgccccatg gctaccgtt gccccactct
2751 gtgagctgat accccattct gggaaactcct ggctctgcac ttcaacctt

Figure 15D

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2801 gctaatatcc acatagaagc taggactaag cccaggaggt tcctctttaa

2851 attaaaaaaaaaaaaaaaaaa

Figure 15E

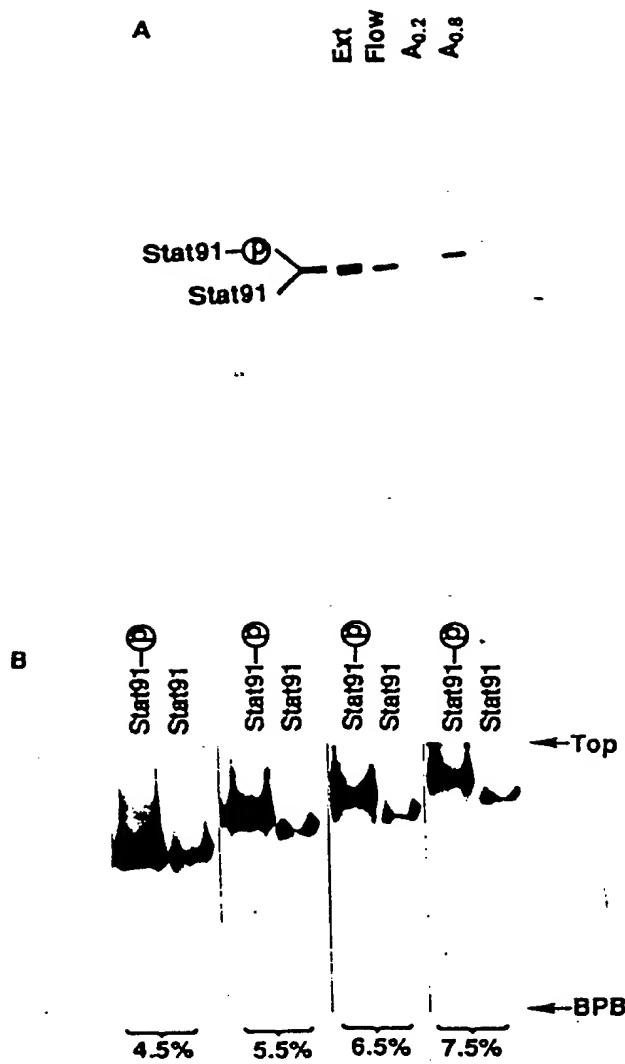


Figure 16A, B

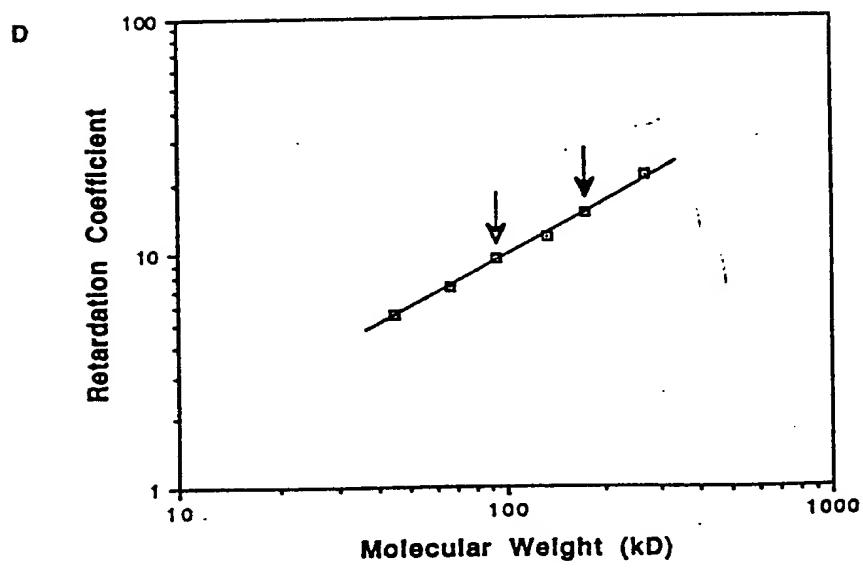
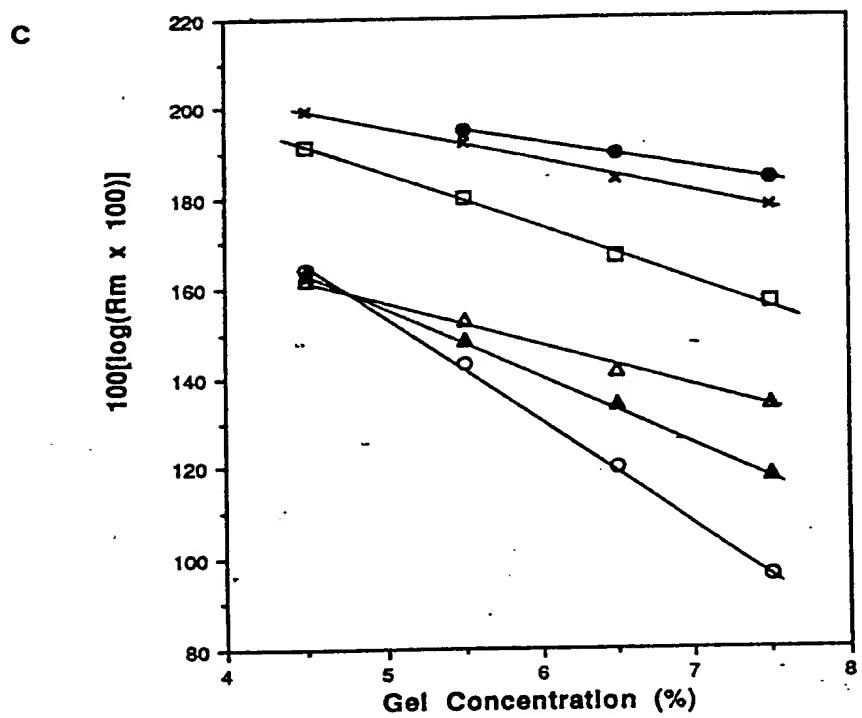


Figure 16C,D

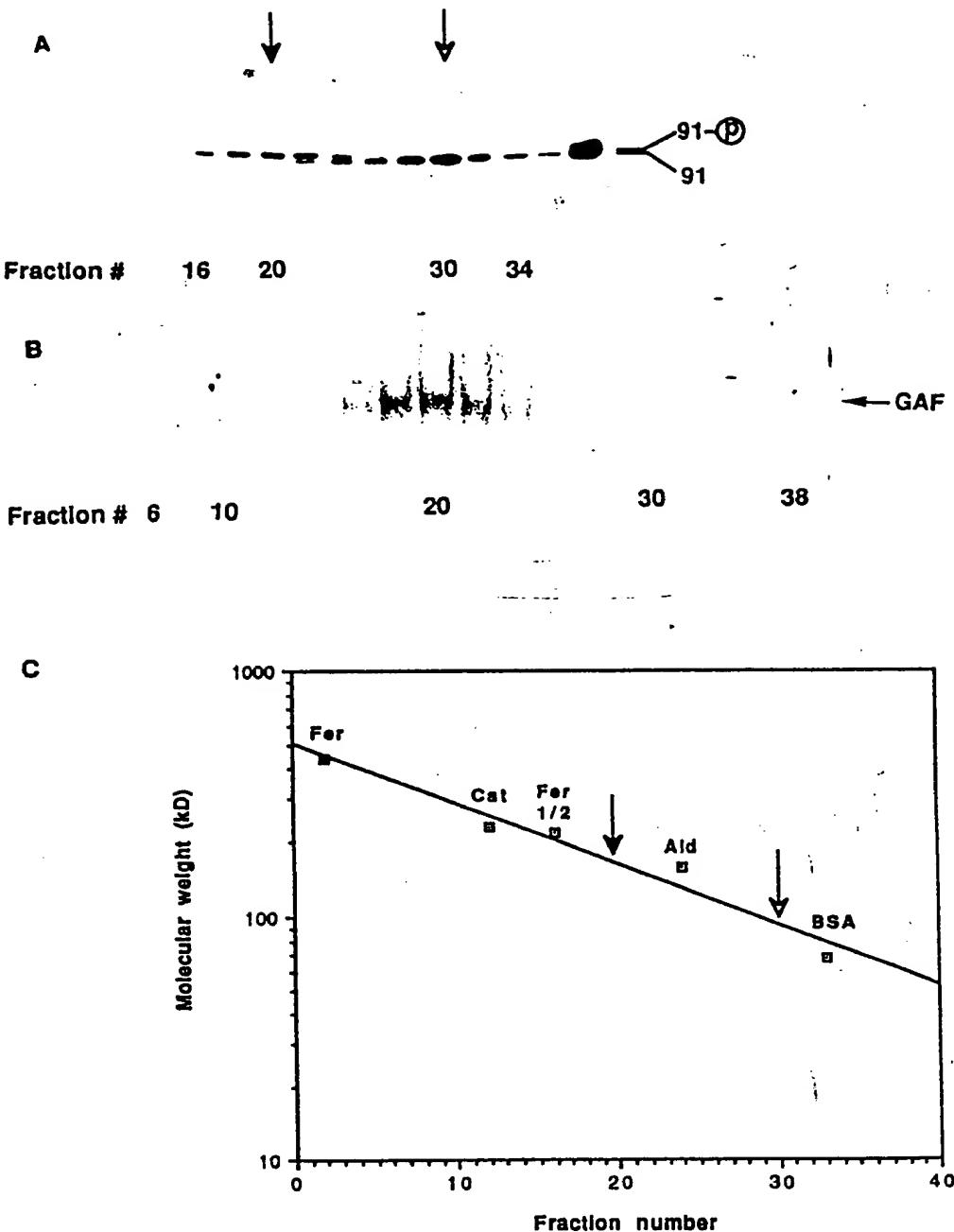


Figure 17

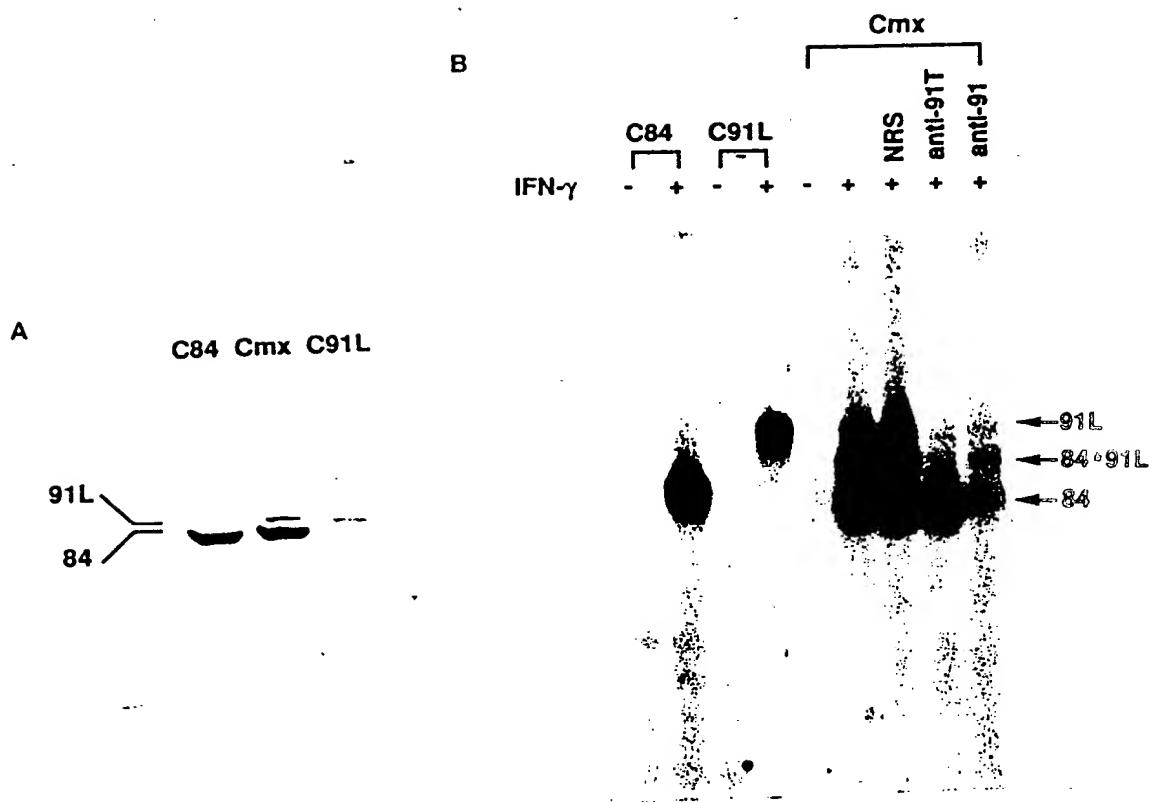


Figure 18

Figure 19

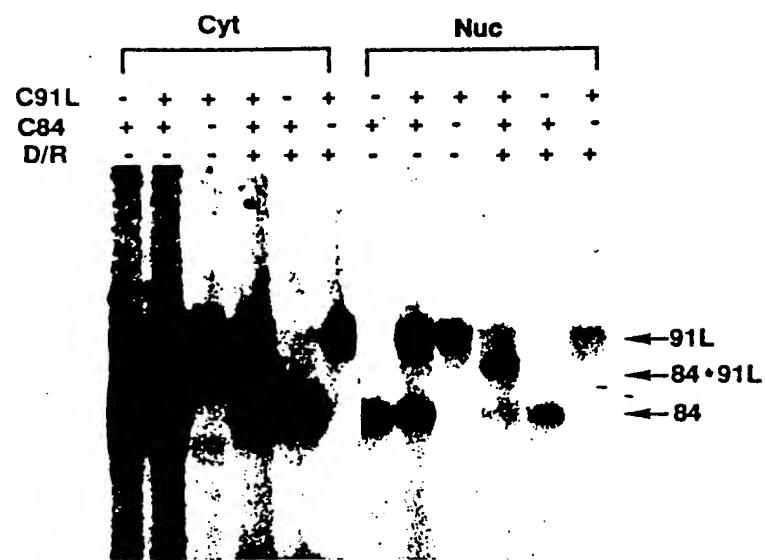


Figure 20

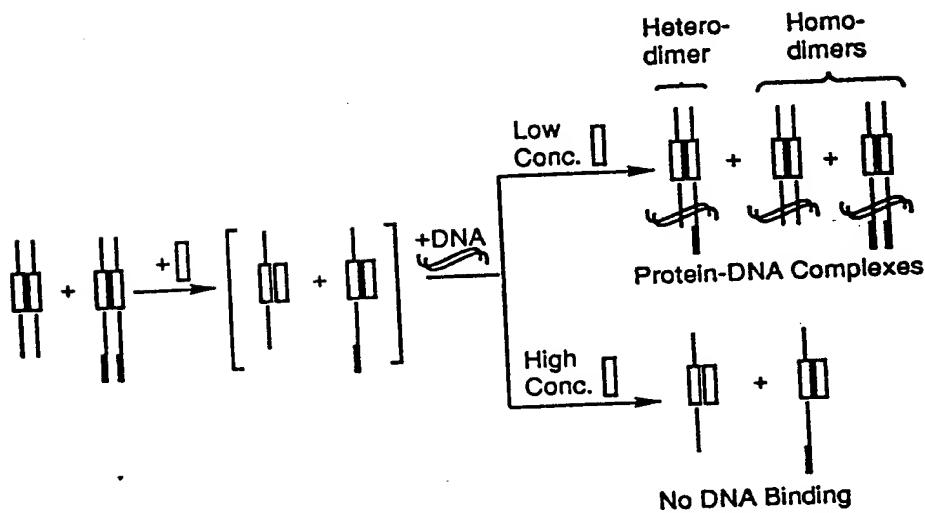


Figure 19, 20

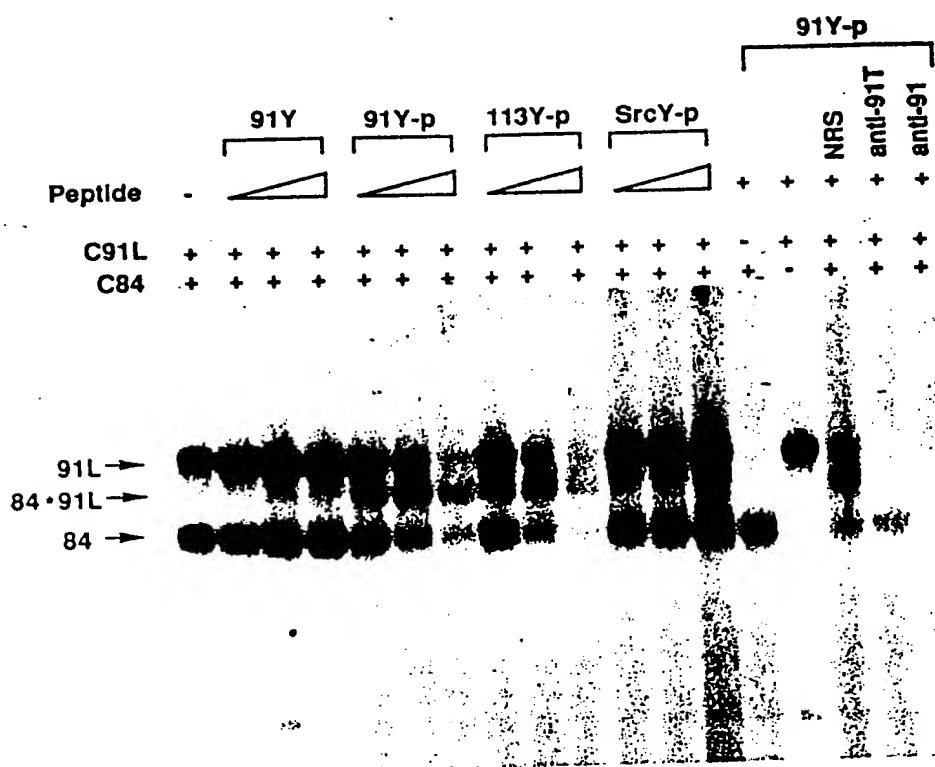


Figure 21

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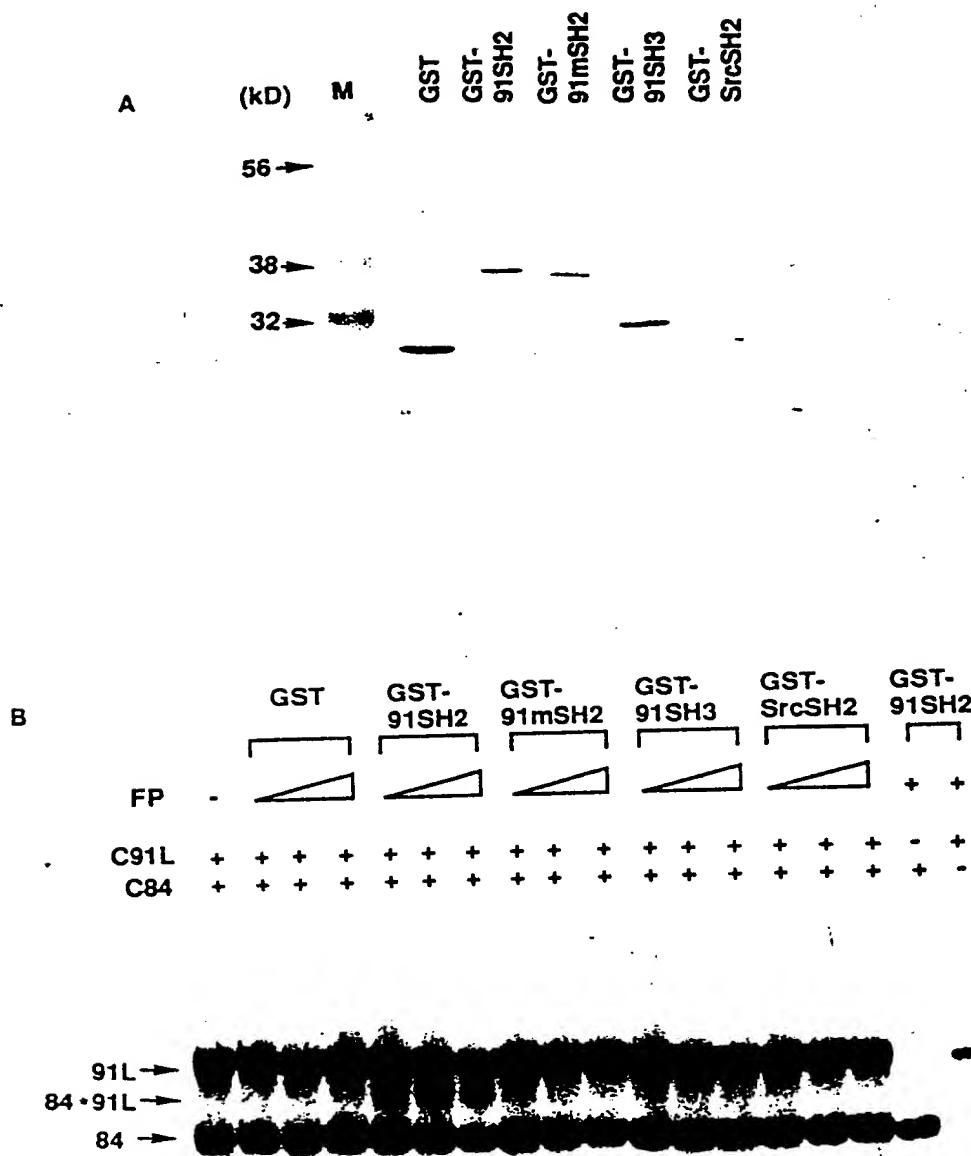


Figure 22

	βA1	αA2	βB5
stat91 (569)	LLPL WND GRCIMGPISKERERALLK DQOP	G TFLLRFS ESSREG AITFWVER	(619)
src (145)	AEE WYF GK1	TRRESERLLL NPENPRG TFLVRES ETTK	G AYCLSVSD (188)
lck (127)	WFF KNL	SRKDAERQLL APGNTHG SFLIRES ESTA	G SFSLSVRD (168)
abl (141)	EKHS WYH GPV	SRNAAEVLLS SGIN G SFLVRES DRRP	G QRISISLRY (184)
p85aN (330)	QDAE WYW GDI	SREEVNEKLR DTAD G TFLVRDA STKMH G DYTTLRLK	(374)

SCR'S	XXX	XXXXXXXXXX	XXXXX	XXX	XXXXXX
Name	[--] [-] [-----]	[-----] [-] [-----]	[-----] [-] [-----]	[-----] [-] [-----]	[-----]
	NA	βA	AA	αA	AB
				βB	BC
				βC	βC

	βD6	
stat91 (620)	S Q N GGEPDFHAVEPYTKKELSAVTFP IIRNYKV MAAENIPENPL (664)	
src (189)	F FD NAK GL	NVKHYKI RKLDS G (210)
lck (169)	D FD QNQ GE	VVKHYKI RNLDN G (189)
abl (185)	E E G	RVYHYRI NTA SD G (200)
p85aN (375)	GG	NNKLIKI FHR D G (388)

SCR'S	XXXXXXXXX X	X
Name	[-----]	[-----] [-] [-----]
	CD	βD
		βD'
		DE

	αB9	
stat91 (665)	KYLY P NID K KDHAFGKYYSRP PK EA PEP M	ELD GPKGTGYIKT (704)
src (211)	GFYI TSR TQP S SLQQQLVAYYSKH AD GL CH	RLT NVC PTS (248)
lck (190)	GFYI SPR ITP P GLHDLVRHYTNA SD GL CT	RLS RPC QTQ (227)
abl (201)	KLYV SSE SRF N TLAELVHHHSTV AD GL IT	TLH YPA PKR (238)
p85aN (389)	KYGF SDP LTP N SVVELINHYRHE S LA QYN PKLDV KL LYP	(427)

SCR'S	XXX	XXXXXXXXXXXX
Name	[--] [-] [-]	[-----] [-----] [-] [-]
	βE	EF
	βF	
		αB
		BG
		βG
		GΩ

Figure 23